## **Ensemble Learning**

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August 1, 2025

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1. Introduction to Ensemble Learning

Use multiple models for prediction. Most winning entries of Kaggle competitions use ensemble learning.

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#### **Example:**

Classifier 1 - Good

Classifier 2 - Good

Classifier 3 - Bad

Using Majority Voting, we predict Good.

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#### **Example:**

Regressor 1 - 20

Regressor 2 - 30

Regressor 3 - 30

Using averaging, we predict  $\frac{80}{3}$ 

Based on Ensemble methods in ML by Dietterich Three reasons why ensembles make sense:

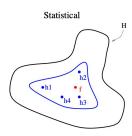
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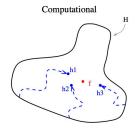
E.g., we can learn many decision trees for the same data giving the same accuracy.



2) Computational: Even if data is sufficient, some classifiers/regressors can get stuck in local optima or apply greedy strategies. Computationally learning the "best" hypothesis can be non-trivial.

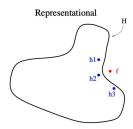
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E.g., decision trees employ greedy criteria

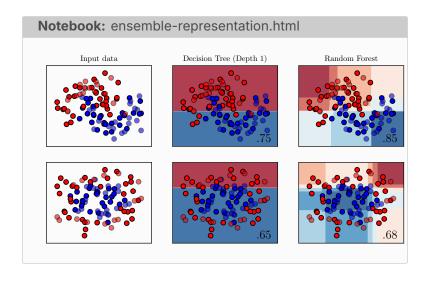


3) Representational: Some classifiers/regressors cannot learn the true form or representation.

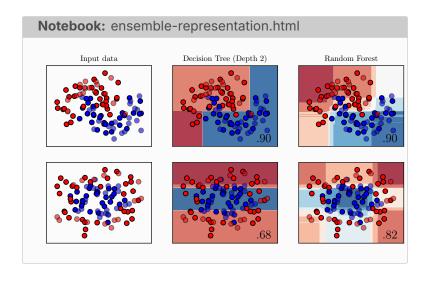
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# Intuition for Ensemble Methods from Quantitative Perspective

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Error Probability of each model =  $\varepsilon$  = 0.3

$$\begin{array}{l} \textit{Pr}(\text{ensemble being wrong}) = \\ {}^{3}\textit{\textbf{C}}_{2}(\varepsilon^{2})(1-\varepsilon)^{3-2} + {}^{3}\textit{\textbf{C}}_{3}(\varepsilon^{3})(1-\varepsilon)^{3-3} \\ = 0.19 < 0.3 \end{array}$$

### Some calculations

Number of Models	Ensemble Error	Individual Error
1	0.3	0.3
3	0.216	0.3
5	0.163	0.3

#### Some calculations

Number of Models	Ensemble Error	Individual Error
1	0.6	0.6
3	0.648	0.6
5	0.683	0.6

Where does ensemble learning not work well?

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- The base model is bad.
- All models give similar prediction or the models are highly correlated.

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Key idea: Reduce Variance

How to learn different classifiers while feeding in the same data?

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Think about cross-validation!

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Key idea: Reduce Variance

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Think about cross-validation!

We will create multiple datasets from our single dataset using "sampling with replacement".

Consider our dataset has n samples:  $D_1, D_2, D_3, \ldots, D_n$ . For each model in the ensemble, we create a new dataset of size n by sampling uniformly with replacement.

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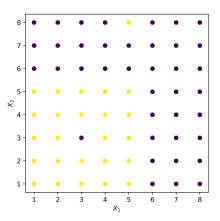
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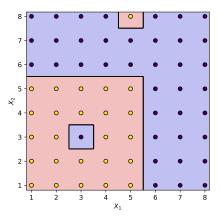
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We can train the same classifier/models on each of these different "Bagging Rounds".

Consider the dataset below. Points (3,3) and (5,8) are anomalies.

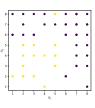


Decision Boundary for decision tree with depth 6.



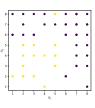
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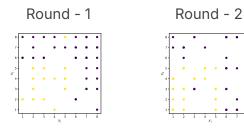
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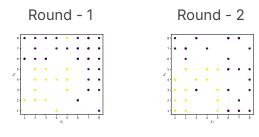


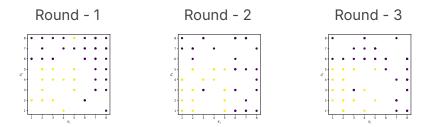
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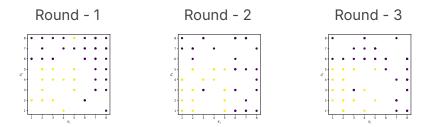
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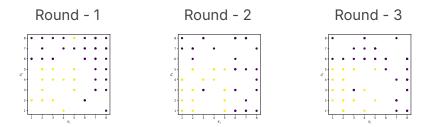


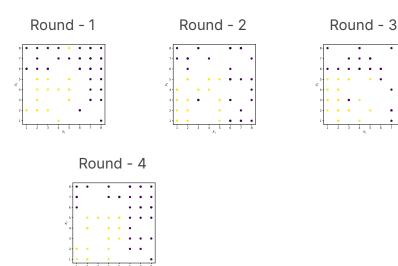


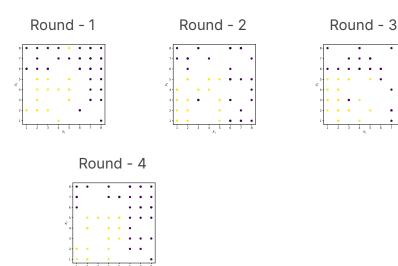


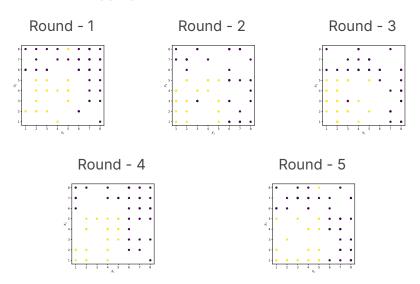




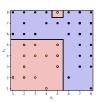






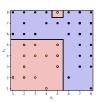


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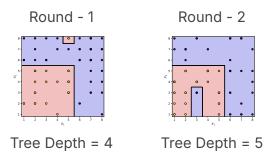


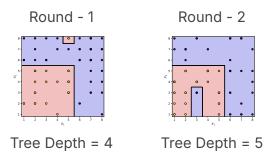
Tree Depth = 4

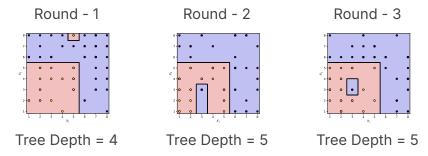
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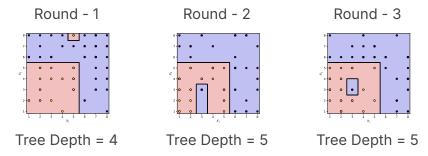


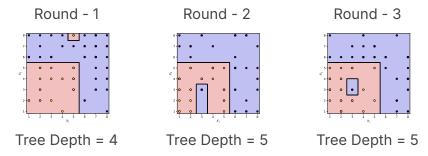
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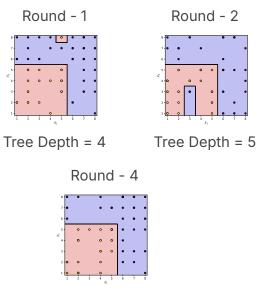


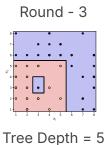


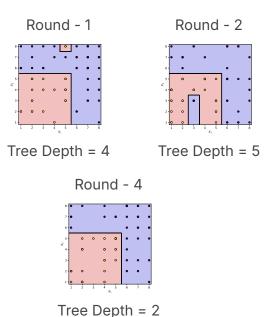


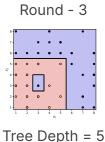


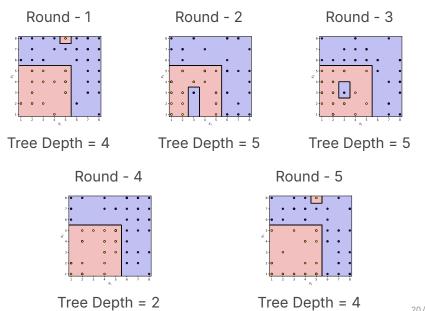






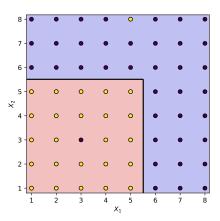






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Using majority voting to combine all predictions, we get the decision boundary below.



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 We take "strong" learners and combine them to reduce variance.

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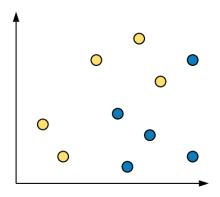
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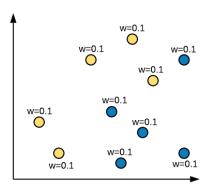
# Boosting

- We take "weak" learners and combine them to reduce bias.
- All learners are incrementally built.
- Incremental building: Incrementally try to classify "harder" samples correctly.



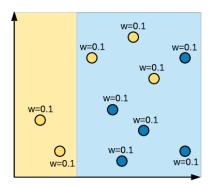
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1. Initialize weights of data samples:  $w_i = \frac{1}{N}$ 

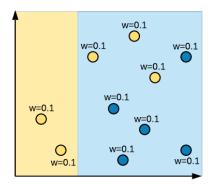


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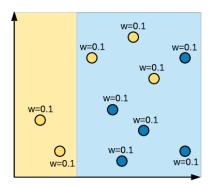
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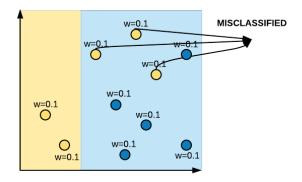


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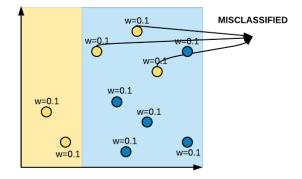


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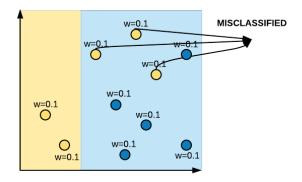
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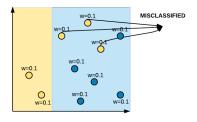
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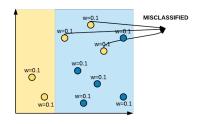
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$$err_1 = \frac{0.3}{1}$$

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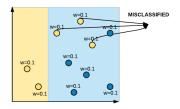
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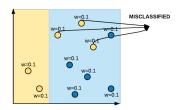
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$$\begin{aligned} &\textit{err}_1 = \frac{0.3}{1} \\ &\alpha_1 = \frac{1}{2} \textit{log}\left(\frac{1-0.3}{0.3}\right) = 0.42 \end{aligned}$$

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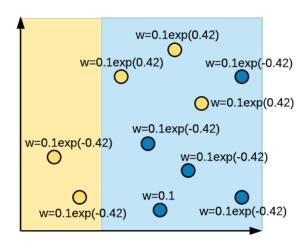
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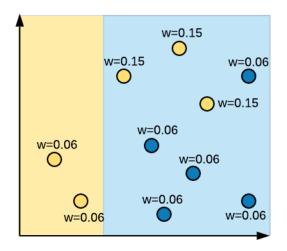
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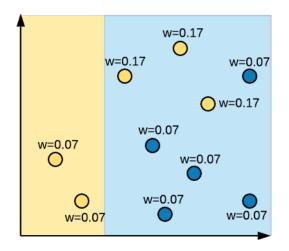
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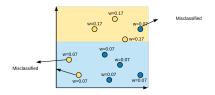
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$$\begin{aligned} \textit{err}_2 &= \frac{0.21}{1} \\ \alpha_2 &= \frac{1}{2} \textit{log} \left( \frac{1 - 0.21}{0.21} \right) = 0.66 \end{aligned}$$

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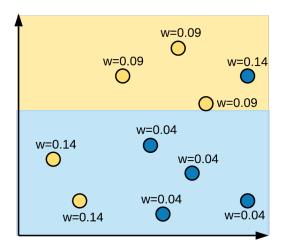
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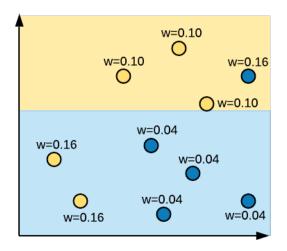
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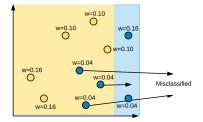
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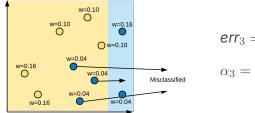
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$$err_3 = \frac{0.12}{1}$$

$$\alpha_3 = \frac{1}{2}log\left(\frac{1 - 0.12}{0.12}\right) = 0.99$$

Intuitively, after each iteration, importance of wrongly classified samples is increased by increasing their weights and importance of correctly classified samples is decreased by decreasing their weights.

#### **Testing**

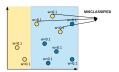
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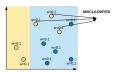
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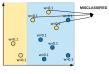
- For each sample x, compute the prediction of each classifier  $h_m(x)$ .
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- SIGN( $\alpha_1 h_1(x) + \alpha_2 h_2(x) + ... + \alpha_M h_M(x)$ )



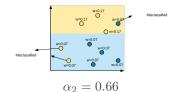
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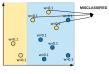


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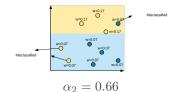


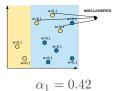
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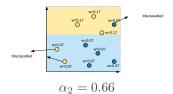


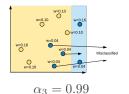


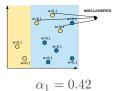
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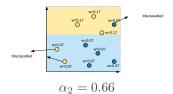


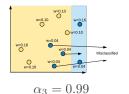


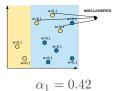


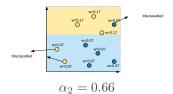


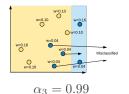


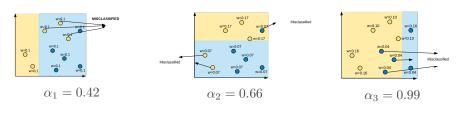


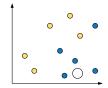


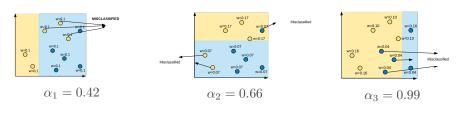


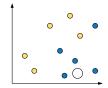


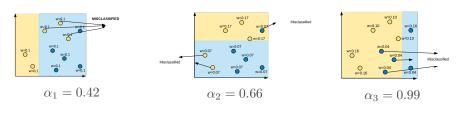


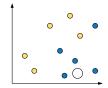




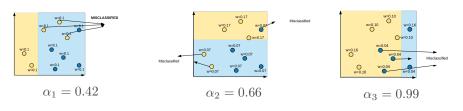


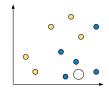






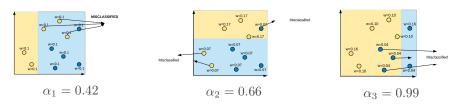
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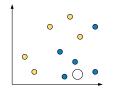




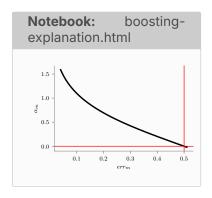
Let us say, yellow class is +1 and blue class is -1

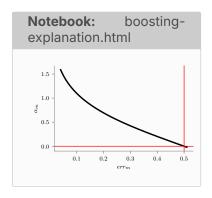
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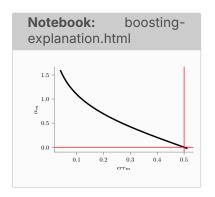


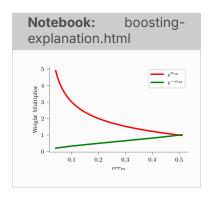


Let us say, yellow class is +1 and blue class is -1 Prediction = SIGN(0.42\*-1 + 0.66\*-1 + 0.99\*+1) = Negative = blue









## **ADABoost for regresion**

# From Paper: Improving Regressors using Boosting Techniques

Our problem will be that the modeling error is also nonzero because we have to determine the model in the presence of noise. Since we don't know the probability distributions, we approximate the expectation of the ME and PE using the sample ME (if the truth is known) and sample PE and then average over multiple experiments.

In the following discussion, we detail both bagging and boosting. We then discuss how to build trees which are the basic building blocks of our regression machines and use these ensembles on some standard test functions.

#### 2. BAGGING

The following is a paraphrase of Breiman (1996b) with some difference in notation. Suppose we pick with replacement N1 examples from the training set of size  $N_{+}$  and call the k'th set of observations  $O_{+}$ . Based on these observations, we form a predictor  $v^{(p)}(\mathbf{x}, O_1)$ . Because we are sampling with replacement, we may have multiple observations or no observations of a particular training example. Sampling with replacement is sometimes termed bootstrap sampling [Efron and Tibshirani (1993)] and therefore this method is called bootstrap aggregating or bagging for short. The ensemble predictor is formed from the approximation to the expectation over all the observation sets, i.e.  $E_O[y^{(p)}(x,O)]$  by using the average of the outputs of all the predictors. Breiman discusses which algorithms are good candidates for predictors and concludes that the best predictors are unstable, i.e., a small change in the training set  $O_k$  causes a large change in the predictor  $v^{(p)}(\mathbf{x}, O_k)$ . Good candidates are regression trees and neural nets.

#### 3. BOOSTING

In bagging, each training example is equally likely to be picked. In boosting, the probability of a particular example being in the training set of a particular machine depends on the performance of the prior machines on that example. The following is a modification of Adaboost A. [Freund and Schapire (1996a)].

Initially, to each training pattern we assign a weight  $w_i=1$   $i=1,...,N_1$ 

Repeat the following while the average loss  $\overline{L}$  defined

set. Each machine makes a hypothesis: h.:x→v

3. Pass every member of the training set through this machine to obtain a prediction  $v^{(p)}(x_i)$   $i=1,...N_1$ .

4. Calculate a loss for each training sample  $L_i - L \left[ \left[ y_i^{(p)}(\mathbf{x}_i) - y_i \right] \right]$ . The loss L may be of any functional form as long as  $L \in [0,1]$ . If we let

$$D=\sup |y_i^{(p)}(\mathbf{x}_i) - y_i|$$
  $i=1,...,N_1$   
then we have three candidate loss functions:

$$L_i = \frac{\mid y_i^{(p)}(x_i) - y_i \mid}{D}$$
 (linear)

$$L_i = \frac{\mid y_i^{(p)}(\boldsymbol{x}_i) - y_i \mid^2}{D^2} \qquad (square~law)$$

$$L_{i} = 1 - exp \left[ \frac{- \mid y_{i}^{(p)}(\mathbf{x}_{i}) - y_{i} \mid}{D} \right] \qquad (exponential)$$

5. Calculate an average loss:  $\overline{L} = \sum_{i=1}^{N_1} L_i p_i$ 

6. Form  $\beta = \frac{\overline{L}}{1 - \overline{L}}$ ,  $\beta$  is a measure of confidence in the predictor. Low  $\beta$  means high confidence in the prediction

7. Update the weights: w<sub>1</sub>→w<sub>1</sub>B\*\*[1-L<sub>1</sub>], where \*\* indicates exponentiation. The smaller the loss, the more the weight is reduced making the probability smaller that this pattern will be picked as a member of the training set for the next machine in the ensemble.

 For a particular input x<sub>t</sub>, each of the T machines makes a prediction h<sub>t</sub>, t=1,...,T. Obtain the cumulative prediction h<sub>t</sub> using the T predictors:

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- We have two types of bagging: bootstrap (on data) and random subspace (of features).
- As features are randomly selected, we learn decorrelated trees and helps in reducing variance.

There are 3 parameters while training a random forest number of trees, number of features (m), maximum depth.

#### Training Algorithm

• For  $i^{th}$  tree ( $i \in \{1 \cdots N\}$ ), select n samples from total N samples with replacement.

#### Learning Decision Tree (for RF)

There are 3 parameters while training a random forest number of trees, number of features (m), maximum depth.

#### Training Algorithm

- For  $i^{th}$  tree ( $i \in \{1 \cdots N\}$ ), select n samples from total N samples with replacement.
- Learn Decision Tree on selected samples for i<sup>th</sup> round.

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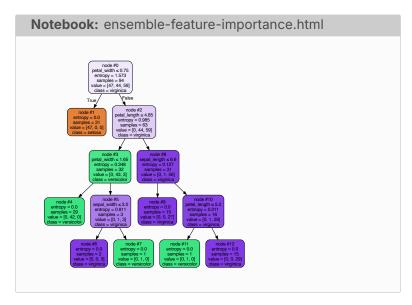
#### Learning Decision Tree (for RF)

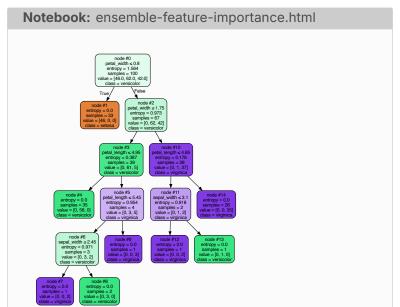
 For each split, select m features from total available M features and train a decision tree on selected features

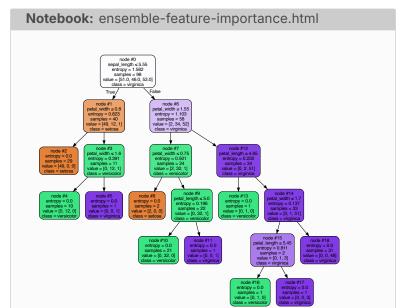
### **Dataset**

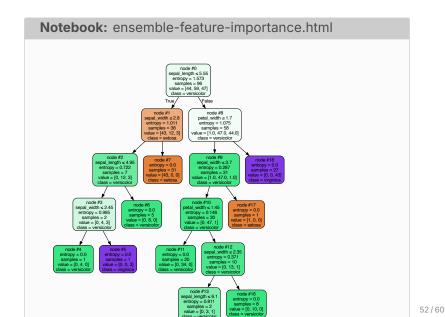
	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa
145	6.7	3.0	5.2	2.3	virginica
146	6.3	2.5	5.0	1.9	virginica
147	6.5	3.0	5.2	2.0	virginica
148	6.2	3.4	5.4	2.3	virginica
149	5.9	3.0	5.1	1.8	virginica

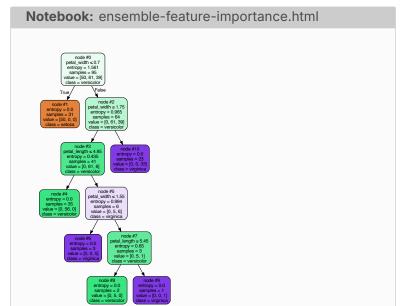
150 rows × 5 columns

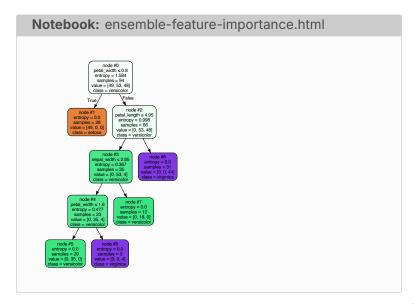


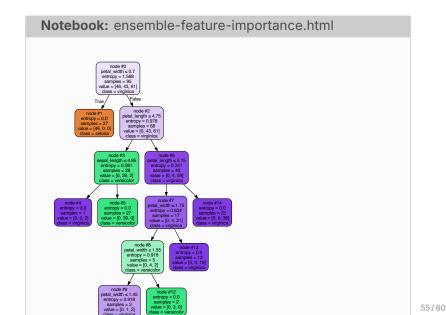


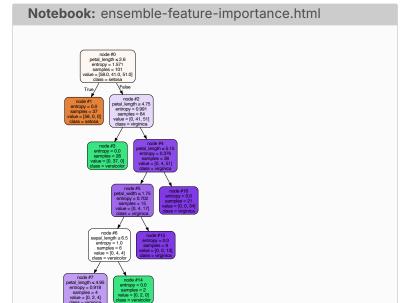


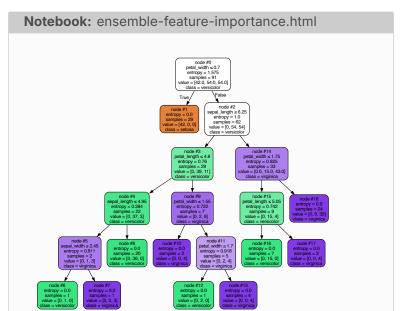


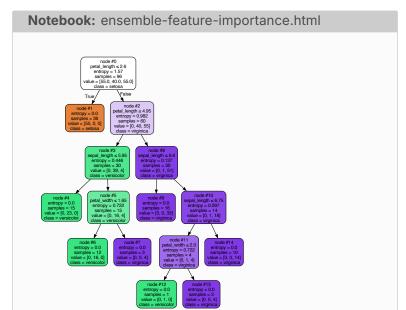




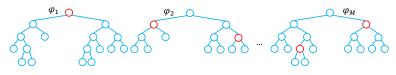








# Feature Importance<sup>1</sup>



Importance of variable  $X_j$  for an ensemble of M trees  $\varphi_m$  is:

$$Imp(X_j) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t \in \varphi_m} 1(j_t = j) \Big[ p(t) \Delta i(t) \Big],$$

where  $j_t$  denotes the variable used at node t,  $p(t) = N_t/N$  and  $\Delta i(t)$  is the impurity reduction at node t:

$$\Delta i(t) = i(t) - \frac{N_{t_L}}{N_t} i(t_L) - \frac{N_{t_r}}{N_t} i(t_R)$$

<sup>&</sup>lt;sup>1</sup>Slide Courtesy Gilles Louppe

# **Computed Feature Importance**

